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Result
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Scoring table:
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                          Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
  167.5
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1: /SIDS5/gcgdata/g
2: /SIDS5/gcgdata/g
3: /SIDS5/gcgdata/g
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5: /SIDS5/gcgdata/g
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Copyright (c) 1993 - 2000 Comp
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AAW37881
AAY30149
AAY30146
AAE02246
AAE02245
AAM79969
AAE02242
AAE02242
AAB69070
AAW19540
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(without alignments)
147.946 Million cell updates/sec
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Human transport as
Domestic mite Htll
Human protein SEQ
Human protein SEQ
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Domestic mite Btll
Human male enhance
Male-enhanced anti
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144.5	144.5	145	145.5	145.5	145.5	146	146	146	147.5	147.5	147.5	147.5	148.5	148.5	149	149.5	150	151	151	151.5	151.5	151.5	151.5	152	152	152	153.5	154	154.5	155	156	156	156
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AAW41586	AAW46823	AAM25750	ABG01716	AAM78854	ABG01723	AAM79504	AAG46981	AAG46982	AAM40883	AAM39097	ABG06505	ABB59344	AAM79864	AAM78880	AAB29659	AAY15457	AAM78520	AAB21227	AAW63043	AAM40467	AAM38681	AAB08316	ABG03671	ABB71125	ABB62094	ABB59948	ABB58673	AAB96721	ABG05140	ABB61289	979	39	AAR72826
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## ALIGNMENTS

RESULT AAW37881

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AAW37881 standard; Protein; 469

AA

28-AUG-1998 (first entry)

AAW37881;

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WPI; 1998-193616/17.
                                           04-SEP-1996;
                                                          06-AUG-1997;
                                                                                                                                                                                        BRCAl modulator
             Ligenfelter C, Polakis P,
                                                                         12-MAR-1998
                                                                                                              Domain
                                                                                                                                     Key
                                                                                                                                                                BRCA1 modulator protein; 091-21A31; breast cancer antigen tumour suppressor protein; diagnosis; therapy; human.
                           (ONYX-) ONYX PHARM INC
                                                                                        WO9810066-A1
                                                                                                                                                    Homo sapiens
                                                                                                                            Domain
                                            96US-0025601
                                                          97WO-US13944
                                                                                                                                                                                        protein 091-21A31.
                                                                                                     /note= "zinc finger motif"
229..255
/note= "leucine zipper motif"
                                                                                                                                   Location/Qualifiers
             Rubinfeld B,
             Vuong TT;
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AAY30149
ID AAY3
XX AAY3
AC AAY3
XX 27-0
DT 27-0
XX Modu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This polypeptide comprises a 53 kDa BRCA1 modulator protein that binds to the tumour suppressor gene product BRC1, and which is characterised by a zinc finger domain and a leucine zipper motif. Its amino acid sequence was deduced from the nucleotide sequence of a cDNA clone (see AAV29062), designated 091-21A31 (ATCC 98141), isolated from a HeLa cell cDNA library using a yeast two-hybrid assay. 3 cDNA clones (see also AAV29063-64) coding for BRCA1 modulator proteins (see AAW39881-83) have been characterised. Vectors and host cells comprising the isolated nucleic acid sequences are claimed, as well as a process for producing BRCA1 modulator protein by culturing these host cells. BRCA1 modulator proteins and nucleic acids can be used to diagnose diseases involving unwanted cell growth, e.g. breast cancer, and to identify compounds that alter BRCA1 interaction with BRCA1 modulators for the treatment of such
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Breast cancer antigen 1 modulator protein - useful for diagnosing diseases involving unwanted cell growth, e.g. breast cancer, and producing therapeutics for treatment of such diseases
                                                                                                                                                                              Modulator protein; BRCA1; tumour suppressor protein; ovarian cancer; cell growth; cell proliferation.
               07-SEP-1999
                                          US5948643-A
                                                                                 Region
                                                                                                              Region
                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                         Amino acid
                                                                                                                                                                                                                                                      27-OCT-1999
                                                                                                                                                                                                                                                                                AAY30149;
                                                                                                                                                                                                                                                                                                           AAY30149 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    217;
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                                                                                                                                                                                                                        sequence of a BRCA1 modulator protein
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                                                                                /note= "zinc finger motif"
230..255
                                                                                                                           Location/Qualifiers
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                                                                    "leucine zipper motif"
                                                                                                                                                                                                                                                                                                          469
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Pred. No. 7.1e-79;
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                                                                                                                                                                                                breast cancer;
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al Similarity 217; Conserv

Conservative

0;

0

Gaps

0;

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Query Match
Best Local S
Matches 217
                                                                                                        The present sequence represents a modulator protein, that binds to and modulate the activity of the BRCA1 gene product (BRCA1). The BRCA1 protein has been characterized as a tumour suppressor protein. Alterations in the amino acid sequence of BRCA1 causes breast and ovarian cancers by removing the controls on cell growth and proliferation. Research has shown that different regions on the BRCA1 molecule have different effects on cell growth and tumour suppression (e.g. full length truncated BRCA1 has no effect on breast cancer cell growth but will inhibit ovarian cancer cell growth). It has been suggested that different bost cell factors (e.g. proteins) interact with different regions of the BRCA1 to control its function. The identification of these proteins (e.g. BRCA1MP) will facilitate the development of novel diagnostic methods and new therapeutics for identifying and treating cancers caused by changes in the expression or activity of BRCA1.
                                                                                                                                                                                                                                                                                                                                                                                      Modulator proteins that bind BRCA1 tumour suppressor gene ovarian and breast cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lingenfelter
                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                         Example 1; Fig 1; 35pp; English
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               98.1%;
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Score 1046; D
Pred. No. 7.1e
0; Mismatches
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           ; DB 20;
7.1e-79;
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for
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                                                                                               ATVVSLQQALGKAEMLCSTLKKQMKYLEQQQDETKQAQEEAGRLRSKMKTMEQIELLLQS
        LQTVYSELDQAKLELKSAQKDLQSADKEIMSLKKKLTMLQ
lqtvyseldqaklelksaqkdlqsadkeimslkkkltmlq
                                                                                                115
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RESULT AAY31646 w

밁 δÃ Вb QY B QΥ В Qy

AAY31646 standard; Protein; 962

A

AAY31646;

02-NOV-1999 (first entry)

Human transport-associated protein-8 (TRANP-8).

Transport-associated vesicle trafficking; hypercholesterolaemia; diagnosis; treatment. protein; TRANP; nuclear |
cancer; cystic fibrosis; nuclear pore; pore; nuclear transport; multidrug resistance;

Homo sapiens

Key

Modified-site

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This sequence represents human transport-associated protein-8 (TRANP-8). The DNA sequence was first identified in a human colon tissue cDNA library. The full-length cDNA was derived from a series of
                                                                  WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                           Modified-site
                                                                               Hillman
                                                                                                                              05-FEB-1999;
                                                                                                                                                         WO9941373-A2
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                          Claim 1; Page 74-77; 87pp; English
                                         cancer
                                             Human TRANP coding sequences, used
                                                                                     Au-Young
                                                                                                   (INCY-) INCYTE PHARM INC
                                                                                                                 11-FEB-1998;
                                                                                                                                           19-AUG-1999
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DB; AAZ11738.
                                                                               JL,
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Lal P,
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902
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reduced
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                                                                            Mite; immunogenic protein; Bt allergen; therapy; atopic dermatitis; immediate hypersensitivity; systemic anaphylaxis; allergic rhinitis asthma; antiallergic; antiinflammatory; immunosuppressive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                            Domestic mite Btll allergen
                                                                                                                                                                                                       31-JUL-2001
                                                                                                                                                                                                                                                                                     AAE02246 standard; Protein; 1017
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                                                                                                                                                            polymorphic variant
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Pred. No. 1.5e-05;
2; Mismatches 85;
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Blomia tropicalis

Location/Qualifiers

Misc-difference

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                                                                                                                                                              Query Match
Best Local :
                                                                                                                                                                                         The present invention relates to immunogenic proteins, referred as Bt allergen, is derived from domestic mite, Blomia tropicalis. The specific Bt allergens of the invention includes Bt11, Bt10, Bt5 and BtA2. The immunogenic protein is useful for preventing, reducing or ameliorating blomia tropicalis hypersensitivity condition such as atopic dermatitis, immediate hypersensitivity, systemic anaphylaxis, allergic rhinitis or asthma and for modulating an immune response directed to Bt allergen in a subject. The Bt allergens are also useful for detecting antibody directed to all or a part of Bt allergen in a biological sample from a subject. Antibodies to Bt allergens are also used as therapeutic or diagnostic agents, to screen Bt immunoassays and as antagonists to inhibit Bt activity under circumstances where temporary hypersensitivity inhibition is required. The present sequence is a protein encoded by matter.
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18-JUL-2000;
18-JUL-2000;
18-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                       Novel immunogenic protein derived from house mite, Blomia tropicalis useful for treating and diagnosing conditions involving induction of immuneresponse to mite, such as allergic asthma, atopic dermatitis,
                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                      Claim 6; Fig
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                                                                                                Local Similarity 22.8 hes 64; Conservative
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                                                                        11 LAQEEENVLD--REFLKNELDN---VRAQLSQ-----KDKEKRDS--
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                                                                                                                                                                                    polymorphic variant.
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71
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56
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22.8%;
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                                                                                            Score 167.5; DB 24; Pred. No. 1.7e-05; """ omatches 74;
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27-APR-2000;
20-JUN-2000;
19-JUL-2000;
01-SEP-2000;
15-SEP-2000;
20-OCT-2000;
                                                                                    The invention relates to polynucleotides (AAK51456-AAK53435) and the encoded polypeptides (AAM78323-AAM80302) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polynucleotides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides have various cytokine-like activities, e.g. stem cell growth factor activity, haematopolesis regulating activity, tissue growth factor activity, immunomodulatory activity and activity inhibin activity and may be useful in the diagnosis and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tang
Zhao
                                                                                                                                                                                                                                                                                                                                                                                                          Nucleic
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                                                                                                                                                                                                                                                                                                                                     Claim
                                                                                                                                                                                                                                                                                                                                                                                    useful
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Wang D,
Yang Y,
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treatment of inflammation

of cancer,

leukaemia, nervous system disorders, arthritis and

Records for SEQ ID NO 2110 (AAK52581), 0020) are omitted as the relevant pages

the relevant pages

2111 from

the

sequence listing

(AAK52582) and 3666

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27-APR-2000;
20-JUN-2000;
                                                                                                                                                                           19-JUL-2000;
01-SEP-2000;
15-SEP-2000;
20-OCT-2000;
                                                                                                                 Tang
Zhao
                                        Nucleic acids encoding polypeptides with cytokine-like useful in diagnosis and gene therapy -
                                                                                                                                                                                                                                                                                                      WO200157190-A2
                                                                                                                                                                                                                                                                                                                                              Human; cytokine; cell proliferation; cell differentiation; gene therapy; vaccine; peptide therapy; stem cell growth factor; haematopoiesis; tissue growth factor; immunomodulatory; cancer; leukaemia; nervous system disorder; arthritis; inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                           AAM79969;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
The invention relates to polynucleotides (AAK51456-AAK53435) and
                      Claim 20;
                                                                        WPI; 2001-476283/51.
N-PSDB; AAK53102.
                                                                                                       Xue
                                                                                                                                               (HYSE-)
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                                                                                                                                                                                                                                                                                                                          Homo sapiens.
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                                                                                                                                               HYSEQ INC.
                                                                                                    Wang D,
Yang Y,
                    Page 397; 6221pp; English
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                                                                                                                                                                  2000US-0560875.
2000US-0598075.
2000US-0620325.
2000US-0654936.
2000US-0653261.
2000US-069325.
2000US-0728422.
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Wang J, Zhang J, Re
Wejhrman T, Goodrich
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Pred. No. 7.6e-06;
0; Mismatches 94;
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RESULT
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Matches 58
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Note: Records for SEQ ID NO 2110 (AAK52581), (AAM80020) are omitted as the relevant pages were missing at the time of publication.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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                                                                                          GEVADK----LRKDLFSSRSKLQTVYSELDQAKLELKSAQKDLQSADKEIMSLKKKLTMLQ
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                                                                                                                                easlldlkehasslassglkkdsrlktleialeqkkeeclkmesqlkkaheaalear-as
                                                                                                                                                                        EEMIRDMGVGQS----
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ilarity 24.2%;
Conservative 5
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                                                                                                                                                                                                                                                                                                                                                              ; Score 167; DB 22;
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50; Mismatches 94;
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AAE02245
                                                   26-OCT-1999;
18-JUL-2000;
18-JUL-2000;
                                                                                                                                                                                                                                                                     Mite; immunogenic protein; Bt allergen; therapy; atopic dermatitis; immediate hypersensitivity; systemic anaphylaxis; allergic rhinitis; asthma; antiallergic; antiinflammatory; immunosuppressive.
                                                                                                                            10-OCT-2000; 2000WO-AU01227
                                                                                                                                                                03-MAY-2001
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                                                                                                                                                                                                                                    Blomia tropicalis
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(UYSI-)
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2000AU-0008844.
2000AU-0008845.
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WPI; 2001-308609/32 N-PSDB; AAD06237.

Chua KY,

Cheong N,

Novel immunogenic protein derived from house mite,

Blomia tropicalis

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Best Local
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            26-OCT-1999;
18-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to immunogenic proteins, referred to as Bt allergens, derived from domestic mite (Blomia tropicalis). The specific Bt allergens of the invention includes Bt11, Bt10, Bt5 and BtA2. The immunogenic protein is useful for preventing, reducing or ameliorating Blomia tropicalis hypersensitivity condition such as atopic dermatitis, immediate hypersensitivity, systemic anaphylaxis, allergic rhinitis or asthma and for modulating an immune response directed to Bt allergen in a subject. The Bt allergens are also useful for detecting antibody directed to all or a part of Bt allergen in a biological sample from a subject. Antibodies to Bt allergens are also used as therapeutic or diagnostic agents, to screen Bt immunoassays and as antagonists to inhibit Bt activity under circumstances where temporary hypersensitivity inhibition is required. The present sequence is Bt11 allergen.
                                                                                                                                                                              Mite; immunogenic protein; Bt allergen; therapy; atopic dermatitis; immediate hypersensitivity; systemic anaphylaxis; allergic rhinitis asthma; antiallergic; antiinflammatory; immunosuppressive.
   18-JUL-2000;
                                                           10-OCT-2000;
                                                                                           03-MAY-2001
                                                                                                                       WO200130817-A1
                                                                                                                                                   Blomia tropicalis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EML-----CSTLKKQMKYLEQQQD----ETKQAQEEAGRLRSKMKTMEQIELLLQS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              875
99SG-0005313.
2000AU-0008842.
2000AU-0008844.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page
                                                                                                                                                                                                                                                                       (first entry)
                                                           2000WO-AU01227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     162-166; 230pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diagnosing conditions involving induction of such as allergic asthma, atopic dermatitis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  60;
                                                                                                                                                                                                                                                                                                                                      878
                                                                                                                                                                                                                                             #7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 166.5;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                      B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .7e-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  74;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22;
                                                                                                                                                                                                                                                                                                                                                                                                                477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                875;
                                                                                                                                                                                               rhinitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  83;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    382
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              322
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Matches
                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   allergen, is derived from domestic mite, Blomia tropicalis. The specific Bt allergens of the invention includes Bt11, Bt10, Bt5 and BtA2. The immunogenic protein is useful for preventing, reducing or ameliorating Blomia tropicalis hypersensitivity condition such as atopic dermatitis, immediate hypersensitivity, systemic anaphylaxis, allergic rhinitis or asthma and for modulating an immune response directed to Bt allergen in a subject. The Bt allergens are also useful for detecting antibody directed to all or a part of Bt allergen in a biological sample from a subject. Antibodies to Bt allergens are also used as therapeutic or diagnostic agents, to screen Bt immunoassays and as antagonists to inhibit Bt activity under circumstances where temporary hypersensitivity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel immunogenic protein derived from house mite, Blomia tropicalis useful for treating and diagnosing conditions involving induction of immuneresponse to mite, such as allergic asthma, atopic dermatitis,
                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to immunogenic proteins, referred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               rhinitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYSI-) UNIV SINGAPORE NAT
                               178
440
                                                                386
                                                                                                  121
                                                                                                                                   326
                                                                                                                                                                                                    266
                                                                                                                                                                                                                                                                       207
                                                                                                                                                                     74
                                                                                                                                                                                                                                     48
                                                                                                                                                                                                                                                                                                      11 LAQEEENVLD--REFLKNELDN---VRAQLSQ-----KDKEKRDS---
                                                                                                                                                                                                                                                                                                                                                      Local
                               RSKLQTVYSELDQAKLELK - - - SAQKDLQSADKEIMSLKKK
                                                                                                  QLPEVEEMIRDMGVGQSAVEQLAVYCVSLKK----EYENLKEARKASGEVADKLRKDLFSS
                                                                                                                                 eygeqleallnkcsalekqkarlqsevevlimdlekatahaqalekrvsqlekinldlks
ksqlndahrriheqeieikrlenereelaaaykeaetlrkq
                                                                  kleevsmll.
                                                                                                                                                                                                  eleslkvqleeesearlelerqltkangdaaswkskyeaelqahvdeveelrrkmaqkis 325
                                                                                                                                                                                                                                                                   lsqenselikevhey-kisldnanhlkgqiaqqledtrhrledeerkrsslenhahtlev 265
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)B; AAD06236.
                                                                                                                                                                      ---EML----
                                                                                                                                                                                                                                    IIDTLRDTLEERNATVVSLQQALGKA-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4.
                                                                                                                                                                                                                                                                                                                                        64;
                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ıs
                                                                                                                                                                                                                                                                                                                                                                                                                         878
                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            230pp;
                                                                                                                                                                   CSTLKKQMKYLEQQQD----ETKQAQEEAGRLRSKMKTMEQIELLLQS 120
                                                               -eqtqkdlrvkiadlqklqheyeklrdqkealarenkkladdlaea
                                                                                                                                                                                                                                                                                                                                                    15.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence is
                                                                                                                                                                                                                                                                                                                                        60;
                                                                                                                                                                                                                                                                                                                                    Score 166.5; DB 2
Pred. No. 1.8e-05;
0; Mismatches 74
                                                                                                                                                                                                                                                                                                                                  74;
                                                                                                                                                                                                                                                                                                                                                                      22;
480
                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                    Length
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                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                      47
                                                                                                                                 385
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73

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AAB69070
                                                                                                   RESULT
                    spermatogenesis
                                         Human male enhanced antigen-2 (MEA-2) protein sequence
                                                        19-APR-2001
                                                                      AAB69070;
                                                                                   AAB69070
                                                                                                  9
                           male
                                                                                    standard;
                           enhanced
                                                       (first
                     disease;
                                                                                    Protein;
                                                       entry)
                           antigen-2;
                     chromosome
                                                                                     1374
                           MEA-2;
                     marker;
                           identification;
                    pancreatic
                    spermatogenesis;
cancer.
                                          SEQ
                                          IJ
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NO:2.

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RESULT 1
AAWA19540
ID AAWA19540
XX AAWA19540
XX AAWA1
XX AAWA1
XX MAle
XX Male
XX Mous
XX Mus
XX Key
FH Key
FH Key
FH Key
FH Miss
XX JP0:
XX JP0:
XX JP0:
XX AAWA1
XX AAWA19540
XX AAWA1954
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Best Local S
Matches 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence represents the human male enhanced antigen-2 (MEA-2). The present invention also described an antibody specific for the MEA-2 protein. The antibody can be used for the identification of a gene causing diseases related to spermatogenesis. The MEA-2 nucleotide sequence is useful as a chromosome marker, and in the detection of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A new protein, human male-enhanced antigen-2, spermatogenesis diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW19540;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 12-15; 21pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-128256/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-NOV-2000
                           07-NOV-1995;
                                                                              13-MAY-1997
                                                                                                                                   JP09121869-A
                                                                                                                                                                                                                 Misc-difference
                                                                                                                                                                                                                                                                                            Mus musculus
                                                                                                                                                                                                                                                                                                                                                    Mouse;
                                                                                                                                                                                                                                                                                                                                                                                                   Male-enhanced antigen-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW19540 standard; Protein; 1325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1176 keqvaaakveaghnrrhfkaaslelsevkkelqakehlvqklqaeaddlqiregkhsqei 1235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1116 rehnsiletalakreadlvqlnlqvqavlqrkeeedrqmkhlvqalqaslekekekvnsl 1175
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JP2000316580-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13 QEEENVLDREFLKNELD-----NVRAQLSQKDKEKRDSQVIIDTLRDTLEERNATVVSL 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             aqfqaelaearaqlqllqkql---deqlskqpvgnqemenlkwevdqkereiqslkqqld 1292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ASGEVADKLRKDLFSSRSKLQTVYSELDQAKLELKSAQKDLQSADKEIMSLKKKL-TMLQ 220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ---QIEL-----LLOSQLPEVEEMIRDMGVGQSAVEQLAVYCVSLKKEYENLKEARK 161
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 QQALGKAEMLCSTLKKQMKYLEQQQDETK------QAQEEAGRLRSKMKTME- 112
                                                                                                                                                                                                                                                                                                                                                 MEA-2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity 25.0
60; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF32508
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1374 AA;
                                                                                                                                                                                                                                                                                               domesticus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                            detecting mutation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99JP-0125196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99JP-0125196
                           95JP-0311638
                                                                                                                                                                                                              Location/Qualifiers 305..320
                                                                                                                                                                                     /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15.4%;
25.0%;
                                                                                                                                                                                  "Not shown in the specification"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 164; DB 22;
Pred. No. 4.9e-05;
46; Mismatches 96;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 1374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   38;
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RESULT
AAW94391
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 57; Conserv
                                                                                                                                                                                                                                                                                                                                                                                               1136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence represents male-enhanced antigen-2 (MEA-2), whi has been derived from a domestic mouse. The polynucleotide encoding the protein can be used for the detection of mutations affecting the MEA-2 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 8;
                                                                                                                                                            Mus
                                                                                                                                                                                Mouse; male enhanced antigen 2; Mea-2; Mus musculus domesticus; spermatogenesis; regulation; contraceptive; sterile; inhibition
                                                                                                                                                                                                                                                                                       AAW94391 standard; Protein; 1325
                                                                                                                                                                                                                                                                                                                                                                                                                                            1086
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                detecting mutation(s)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Male-enhanced antigen Mea-2 gene - especially from mouse, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1997-314229/29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-NOV-1995;
                                                                  04-JUL-1997;
                                                                                        04-JUL-1997;
                                                                                                               26-JAN-1999
                                                                                                                                     JP11018622-A.
                                                                                                                                                                                                                    Mouse
                                                                                                                                                                                                                                           14-APR-1999
                                                                                                                                                                                                                                                                 AAW94391;
                                                                                                                                                                                                                                                                                                                                                1193 nmktllq 1199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N-PSDB; AAT74034.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ITOH-) ITO HAM KK
                                           (ITOH-) ITO HAM KK.
                                                                                                                                                                                                                                                                                                                                                                       215
                                                                                                                                                                                                                                                                                                                                                                                                                     155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  966 rehnsiletalakreadlvqlnlqvqavlqrkeeedrqmkqlvqalqvslekekmevnsl 1025
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13
                                                                                                                                                                                                                                                                                                              11
                                                                                                                                                                                                                                                                                                                                                                                                          NLKEARKASGEVADKLRKDLFSSRSKLQTVYSELDQAKLELKSAQKDLQSADKEIMSLKK
                                                                                                                                                                                                                                                                                                                                                                       KL-TMLQ 220
                                                                                                                                                                                                                                                                                                                                                                                             slkqqldlteqqg---kkelegtqqtlqtikselemvqedlsetqkdkfmlqakvselkn 1192
                                                                                                                                                                                                                                                                                                                                                                                                                                           agfqtelaeartqlq-----llqkkl---deqmsqqptgsqemedlkweldqkereiq 1135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  KQAQEEAGRLRSKMKTMEQIELLLQSQLPEVEEMIRDMGVGQSAVEQLAVYCVSLKKEYE 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       keqmaaarieaghnrrhfkaatlelsevkkelqakehlvqtlqaevdelqiqdgkhsqei 1085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                QQALGKA-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           QEEENVLDREFLKNELD-----NVRAQLSQKDKEKRDSQVIIDTLRDTLEERNATVVSL
                                                                                                                                                                                                                   male
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Page 9-10; 13pp; Japanese
                                                                                                                                                                                                                   enhanced antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                          (first entry)
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                                                                 97JP-0179490
                                                                                        97JP-0179490
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Pred. No. 8.4e-05;
8; Mismatches '90,
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7;

WPI; 1999-160962/14 N-PSDB; AAX04132.

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RESULT 12
AAR72826
ID AAR728
XX AAR728
XX AAR728
XX CALL
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CKW INhibi
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR72826
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                                  24-OCT-1994;
                                                                                         27-APR-1995
                                                                                                                                         WO9511309-A2
                                                                                                                                                                                                                          Region
                                                                                                                                                                                                                                                                                Region
                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                          inhibition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human mitosin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-FEB-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR72826 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 cycle; M
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        KQAQEEAGRLRSKMKTMEQIELLLQSQLPEVEEMIRDMGVGQSAVEQLAVYCVSLKKEYE 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity 23.1
57; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1325 AA;
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                                  94WO-US12162
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1660..18
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23.1%;
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1839
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3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 161; DB 20;
Pred. No. 8.4e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 retinoblastome; mitosis; cell growth;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      90;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 1325;
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AAR72829 is human mitosin. Mitosin is involved in the regulation of the mammalian mitotic cell cycle. Mitosin as with E2F-1 (see AAR72824) interacts with the retinoblastoma protein (the retinoblastoma tumour suppressor gene product). Mitosin is first synthesised at the G1/S boundary, it is then phosphorylated from S through M phase, and during mitosis, is closely associated with the centromeres/kinetochores at the mitotic spindle poles. Mitosin is necessary for a eukaryotic cell to enter the M phase of the mitotic cell cycle and its degradation is necessary for a cell to advance on to the next stage. Mitosin is thus useful for controlling cell growth as overexpression of mitosin prevents a cell from exiting the M phase.
                                                            An anti-mitosin antibody, antibody fragment or a phosphorylated mitosin mutein ( or nucleic acid encoding it) can also be used to inhibit cell division which is particularly useful for the study of the cell cycle. A further use is to control hyperproliferative cells, and so control diseases such as psoriasis and breast cancer. It can also be used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Purified mammalian protein mitosin and agents that bind inhibit its action - used to promote cell growth or to division and/or proliferation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1995-170229/22
N-PSDB; AAQ86851.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lee
Sequence
                                          block gametogenesis of an immature gamete.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (TEXA ) UNIV TEXAS
                                                                                                                                                                                                                                                                                                                                                                                                                                   Fig 8B; 61pp; English
    2482 AA;
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Query Match
Best Local S
Matches 57
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                                         1751 leldvvtirsekenltnelqkeqeriseleiinssfenilqekeqekvqmkeksstamem 1810
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                                                                                                                                                                                                                                                                                           1571 ldlvtlrsekenltkqiqekqgqlseldkllssfkslleekeqaeiqikeesktavemlq 1630
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219
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                                                                                LRKDLFSSRSKLQTVYSELDQ-----AKLEL--KSAQKDLQSADKEIMSLKKK----LTM 218
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220
                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                               -----VEEMIRDMGVGQSAVEQLAVYCVSLKKEYENLKEARKASGEVADK 169
                                                                                                                                                                                                                                                    ----MKYLEQQQD----ETKQAQEEAGRLRSKMKTMEQIELLLQS 120
                                                                                                                                                                                                                                                                                                                                                                                               14.6%;
                                                                                                                                                                                                                                                                                                                                                                              55;
                                                                                                                                                                                                                                                                                                                                                                            Score 156; DB 16;
Pred. No. 0.00047;
5; Mismatches 90;
                                                                                                                                                                                                                                                                                                                                                                                                                 Length 2482;
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                                                                                                                                                                                                                                                                                                                                                                              40;
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7;

AAW23996
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AC AAWZ
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AC AAWZ
XX
DT 28-N
DT 28-N
DT Unne
XX
DE Hume
XX
KW Mitc

AAW23996;

AAW23996 standard; Protein;

2482

Human mitosin amino acid sequence

(first entry)

Mitosin; phosphoprotein; mitotic cell cycle; antibody; analogue; inhibition; M phase; Antagonist; hyperproliferative cell; cancer;

Вb Š В Qy Вp 8 뫄 QY В Q

1811

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1812

RESULT 13

leukaemia; lymphoma;

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Query Match
Best Local Similarity
Matches 57; Conserv
                                                                                                                                          anti-mitosin antibody, a mutant or a non-functional analogue of mitosin can inhibit the mitotic cell cycle by preventing the cells from entering the M phase, and over expression of mitosin or its functional equivalent, would inhibit the cycle by preventing cells from leaving the M phase. Antagonists to this protein can be used to control experproliferative cells in, (e.g. thyroid hyperplasia, Grave's disease, psoriasis, benign prostatic hypertrophy, Li-Fraumeni syndrome, breast cancer; sarcomas and other neoplasms, bladder cancer, colon cancer, lung cancer and various leukaemias and lymphomas). Reintroduction of supplementation of lost mitosin function by introduction of the protein or nucleic acid encoding the protein into a cell can restore defective
                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Domain
Domain
Domain
Domain
                                                                                                     chromosome segregation, which is a marker of progressing malignancy. Malignant proliferation of cells can then be halted. The protein can also be used for the detection and diagnosis of hyperproliferative
                                                                                                                                                                                                                                                                                             This is the amino acid sequence for mitosin, a phosphoprotein necessary for the cell to ente mitosis. The protein's degradatic also necessary for the cell to advance into the next stages of mitosin protein, can be used to control the growth of cells.
                                                                                                                                                                                                                                                                                                                                                                                          New isolated mitosin protein and gene - useful for, e.g. developing products for therapy and diagnosis of hyper-proliferative disorders such as cancers or psoriasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Domain
                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lee
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-OCT-1994;
22-OCT-1993;
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Misc-difference
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bB; AAV09076.
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                                                                   2482
  Conservative
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93US-0141239
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2189
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2303
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1885..1962
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/note= "optinally A or T"
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re= "leucine h
            14.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     chromosome segregation.
                                                                                                                                                                                                                                                                                                                                                                 43pp; English
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"Optionally C or G"
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  55;
Score 156; DB 19;
Pred. No. 0.00047;
5; Mismatches 90
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  90;
                         Length 2482;
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DNA encoding kinetochore protein - uphases of a cell cycle, partic. for
                              WPI; 1996-287116/29
N-PSDB; AAT34578.
                                                                                                               09-DEC-1994;
                                                                                                                                    08-DEC-1995;
                                                                                                                                                                                                                                                             Domain
                                                                                                                                                                                                                                                                                                                                              Domain
                                                                                                                                                                                                                                                                                                                                                                              Key
                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR99795;
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                                                             Rattner JB,
                                                                                (FOXC-) FOX CHASE CANCER CENT (UYTE-) UNIV TECHNOLOGIES INT
                                                                                                                                                         13-JUN-1996
                                                                                                                                                                            WO9617867-A1
                                                                                                                                                                                                                                          Domain
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         220
                                                                                                                                                                                                                                                                                                                                                                                                                     e protein; antibody.
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                                                                                                               94US-0353700
                                                                                                                                    95WO-US16216
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/note= "
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/label= Globular_domain
                                                                                                                                                                                                                                                                         /label=
                                                                                                                                                                                                                                                                                                                                                                             location/Qualifiers
                                                                                                                                                                                                                                                    /label=
                                                                                                                                                                                                                                                                                                                                    'Label=
                                                                                                                                                                                                                                                                                                                                                        'label= Extended_coiled_structure
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                                                                                                                                                                                               "the C-terminal domain is predicted form a proline-rich (10.6%) highly basic (pI 10) globular domain"
                                                                                                                                                                                                                                                                                                      "globular
                                                                                                                                                                                                                               C-terminal_domain
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                                                                                                                                                                                                                                                   Extended_coiled_structure
                                                                                                                                                                                                                                                                                            repeats of 95 amino
                                                                                  INT INC
                                                                                                                                                                                                                                                                                                                                                                                                                              cell cycle; cancer; diagnosis;
- used as a marker for the {\sf G2} and {\sf M} for detection of malignant diseases
                                                                                                                                                                                                                                                                                                       domain
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Best Local S
Matches 57
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                        N-PSDB;
                                                                                                                                                                                                                Drosophila melanogaster
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  New
                                                            Venter JC, Adams M,
                                                                                  (PEKE ) PE
                                                                                                          23-MAR-2000;
11-JUL-2000;
                                                                                                                                              23-MAR-2001; 2001WO-US09231
                                                                                                                                                                     27-SEP-2001
                                                                                                                                                                                           WO200171042-A2
                                                                                                                                                                                                                                          pharmaceutical.
                                                                                                                                                                                                                                                       Drosophila;
                                                                                                                                                                                                                                                                          Drosophila melanogaster polypeptide SEQ ID NO 10659
                                                                                                                                                                                                                                                                                                   26-MAR-2002
                                                                                                                                                                                                                                                                                                                           АВВ61289;
                                                                                                                                                                                                                                                                                                                                                  ABB61289 standard; Protein; 455 AA
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  isolated nucleic acid
                        2001-656860/75
DB; ABL05392.
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                                                                                    CORP NY.
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                                                                                                                                                                                                                                                    developmental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                          2000US-191637P
2000US-0614150
                                                                                                                                                                                                                                                                                                  (first entry)
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                                                            Li
                                                           , DWD
detection reagent for detecting 1000 or more
                                                                                                                                                                                                                                                    biology;
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Pred. No. 0.00066;
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                                                           Myers
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                                                                                                                                                                                                                                                  cell signalling; insecticide;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 3248;
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                                                                                                                                                                                                                                                                                                                                                                              RESULT 16
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Best Local Similarity
Matches 57; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          capable of detecting 1000 or more genes from Drosophila. The invent useful in developmental biology and in elucidating cell signalling cell-cell interactions in higher eukaryotes for the development of insecticides, therapeutics and pharmaceutical drugs. The invention discloses genomic DNA sequences (ABLJ6176-ABL30511), expressed DNA sequences (ABLJ617737-ABB72072).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from at ftp.wipo.int/pub/published_pct_sequences.
                                                                 31-MAR-2000;
23-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   genes from Drosophila and for elucidating cell signalling and cell-cell interactions \boldsymbol{\cdot}
                                                                                                          30-MAR-2001; 2001WO-US08631
                                                                                                                                                                 WO200175067-A2
                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                     Human; chromosome mapping; gene mapping; gene
food supplement; medical imaging; diagnostic;
                                                                                                                                                                                                                                                                 Novel
                                                                                                                                                                                                                                                                                             13-FEB-2002
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                                                                                                                                       11-OCT-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 dadrllkn----eadphvlanwvstlkrelrqceskktelrnvvkvvqndlrkeielkry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        agleqkvqkkdflissyveqigvlksdahvvdglrkenktlksqiqsmegisailaagsa 174
                                                                                                                                                                                                                                                                human diagnostic protein #5131.
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                                                                 2000US-0540217
2000US-0649167
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Pred. No. 7e-05;
58; Mismatches ?
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                                                                                                                                                                                                                     therapy;
genetic 
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Matches
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    27-OCT-2000
                                               FR2792651-A1
                                                                                       Pyrococcus abyssi
                                                                                                                               Hyperthermophilic
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                                                                                                                                                                                                                                                                                                        AAB96721 standard; Protein; 1177
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DB; AAS69327.
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                                                                                                                             hyperthermophilic protein.
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Querellou J,
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J, Saurin
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pharmaceutical.
               Drosophila; developmental biology;
                                                  Drosophila melanogaster polypeptide SEQ
                                                                                             26-MAR-2002
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  1898
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          biology;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    detection reagent for detecting 1000 for elucidating cell signalling and
                                                                                                                                                                                                                                                                                                                                                                                                                          49;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 152; DB 22; Pred. No. 0.00081;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Myers
                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          signalling; insecticide
                                                                                                                                                                                                                                                                                                                                                                                                                        84;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        format directly from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                part of the printed
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 2067;
                                                                                                                          ---AKL--ELKSAQKD
                                                                                                                                                                 eladaheqlnevsaq
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                                                                                                                                                                                                                                                                                                                                                                                                                        64;
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standard;

Protein;

ABG03671;

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                                                                                                                                                                                                                                                                                                                       cc polypeptide (II) sequences. (I) is useful as hybridisation probes, cc polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The care also used in diagnostics as expressed sequence tags cfor identifying expressed genes. (I) is useful in gene therapy techniques (II). (II) is useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful for treating disorders involving aberrant protein expression or biological activity. The polypeptide and polynucleotide sequences have applications in diagnostics, forensics, gene mapping, identification of mutations cresponsible for genetic disorders or other traits to assess biodiversity and to produce other types of data and products dependent on DNA and amino acid sequences. ABG00010-ABG30377 represent novel human and alternative to the content of the invention.
                                                                                                                                                                                     Query Match
Best Local
                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess
                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                             Note: The sequence data for this patent did not appear in the specification, but was obtained in electronic format directly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 20; SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (HYSE-)
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23-AUG-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel human diagnostic protein #3662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-FEB-2002
                                                215
                                                                                                           155
                                                                             71
                                                                                                                                        12
                                                                                                                                                                                                                                                                             ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        invention relates to isolated polynucleotide (I) and
                                           kkvraelerwrkaaseyekevtslqnsfqlrcqqcedqqreeasrlqgeleklrkewnal\\
                 E----
                                                                           GK--AEM----LCSTLKKQMKYL---
                                                                                                      aqwqrfhidtenlreqkdneias ardelh sardem wlvh qaaak vaserdt dia slqeel\\
                                                                                                                                     AQEEENVLDREFLKNELDNVRAQLSQKDKEKRDSQVII-DTLRDTLEERNATVVSLQQAL 70
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DB; AAS67858.
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                                                                                                                                                                                     Similarity
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                                                                                                                                                                        Conservative
                -QIELLLQSQLPEVE---
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2000US-0649167
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mapping; gene mapping; gene therapy;
medical imaging; diagnostic; genetic (
                                                                                                                                                                                   14.2%;
24.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TY
                                                                                                                                                                      43;
                                                                                                                                                                   Score 151.5; DB 2
Pred. No. 0.00014;
13; Mismatches 102
              EMIRDMGVGQSAVEQLAVYCVSLKKEY
                                                                         EQQQDETKQAQEEAGRLRSKMKTM
<del>..</del>
                                                                                                                                                                                                   DB 22;
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                                                                                                                                                                      Indels
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                 153
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                DЪ
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Query Match Best Local Matches

Local

Similarity 21.3 50; Conservative

14.2%; 21.3%;

Score 151.5; DB 21; pred. No. 0.00026; 62; Mismatches 84;

21;

753; 39;

Indels Length

Gaps

6,

63 12

> eelegqieklqaevkgykdennrlkekehknqddllkeketliqqlkeelqeknvtldvq EFLKNELDNVRAQL-SQKD-----KEK-------RDSQVIIDTLRDTLEERNAT----

72

iqhvvegkralseltggvtcykakikeletiletqkvershsakleqdilekesiilkle --- VVSLQQALGKAEMLCSTLKKQMKYLEQQQDETKQAQEEAGRLRSKMKTMEQIELLLQ

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAB08316
                                    The present sequence represents a human M-phase phosphoprotein-1 (MPP1) polypeptide. MPP1 is strongly expressed in mitotic cells, and is synthesised during interphase of the cell cycle. A subset of idiopathic ataxia patients have autoantibodies to the antigen MPP1, and detection of these autoantibodies is useful in classifying and identifying the type of ataxia experienced by the patient. The specification describes a method for detecting the presence of autoantibodies to MPP1 a biological sample from a patient diagnosed with idiopathic ataxia.
                                                                                                                                                                                                  Detecting presence of an antibody associated with ataxia in a biological sample for diagnosing and treating ataxia, involves contacting the sample with a peptide comprising a sequence identical to
                          The detection
                                                                                                                                                            Claim 35; Page 58-60;
                                                                                                                                                                                                                                                        N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A human M-phase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-DEC-2000
Sequence
                                                                                                                                                                                        M-phase
                                                                                                                                                                                                                                                                                                                                                 09-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                            09-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; M-phase phosphoprotein-1; MPP1; mitotic
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                                                                                                                                                                                                                                                                                                                        (UYTE-) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      idiopathic ataxia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               335
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ENLKEARKASGEVADKLRKDLFSSRSKLQTVYSELDQAKLELKSAQKDLQSADKEIMSLK
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                                                                                                                                                                                      phosphoprotein-1
                                                                                                                                                                                                                                                         AAA63952.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             398
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                          method
AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antigen; autoantibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      entry)
                           is
                                                                                                                                                             72pp; English.
                          useful for diagnosing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polypeptide
                          and treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cell; interphase;
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RESULT
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14-SEP-2000;
19-OCT-2000;
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25-APR-2000;
09-JUL-2000;
19-JUL-2000;
                                system, such as peripheral nervous injuries, peripheral neuropathy and localised neuropathies and central nervous system disease, such as Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager Syndrome. Other uses include the utilisation of the activities such as: Immune system suppression, Activinyinhibin activity, chemotactic/chemokinetic activity, haemostatic and thrombolytic activity, cancer diagnosis and therapy, drug screening, assays for receptor activity, arthritis and inflammation, leukaemias and assays for receptor activity, arthritis and inflammation, leukaemias and
                                                                                                                                    The invention relates to human nucleic acids (AAI57798-AAI61369) and the encoded polypeptides (AAM38642-AAM42213) with nootropic, immunosuppressant and cytostatic activity. The polypucleotides are useful in gene therapy. A composition containing a polypeptide or polynucleotide of the invention may be used to treat diseases of the peripheral nervous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; nootropic; immunosuppressant; cytostatic; gene therapy; cancer; peripheral nervous system; neuropathy; central nervous system; CNS; Alzheimer's; Parkinson's disease; Huntington's disease; haemostatic; amyotrophic lateral screensis; Shy-Drager Syndrome; chemotactic; chemokinetic; thrombolicia.
                       assays for receptor activity, arthritis {\tt C.N.S} disorders.
                                                                                                                                                                                                                                                                                                                           Zhao
                                                                                                                                                                                                                                                                                                                                       Wang
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specification
          Note: The sequence
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                                                                                                                                                                                                                                                                                                                         J,
QA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sapiens.
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)B; AAI57837.
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                                                                                                                                                                                                                                              nucleic acids and as central nervous
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                                                                                                                                                                                                                                                                                                                          Liu C
Wang Z
Zhou
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2000US-0620312.
2000US-0653450.
2000US-0662191.
2000US-0693036.
2000US-0727344.
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2000US-0552317.
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                                                                                                                                                                                                                   ID NO 1826; 10078pp; English.
                                                                                                                                                                                                                                                                                                                                      Asundi V,
Wehrman T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Protein;
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            data
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Xu C, Xue AJ,
R, Drmanac RT;
          patent
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            did
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          form
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25-APR-2000;
09-JUL-2000;
19-JUL-2000;
19-JUL-2000;
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                                                                                        Wang
                                                                                                Tang
                                                                                                                                                                                                                                                                                                          Human: nootropic; immunosuppressant; cytostatic; gene therapy; cancer; peripheral nervous system; neuropathy; central nervous system; CNS; Alzhelmer's; Parkinson's disease; Huntington's disease; haemostatic; amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                          such
                                                                               Zhao
                                                                                                                                                                                                                                                                                                                                                         Human polypeptide
                                                                                                                                                                                                                                                                                                                                                                                             AAM40467;
                                                                                                                                                                                                                                                                                                                                                                                                               AAM40467
                                                                                                                                   29 - NOV -
                                                                                                                                           14-SEP-2000;
19-OCT-2000;
                                                                                                                                                                                                                                                      WO200153312-A1
                                                                                                                                                                                                                                                                                                   chemokinetic;
                                                                                                                                                                                                                                                                                                                                                                           22-OCT-2001
                                                                                                                (HYSE-)
                                                                                                                                                                                                                   26-DEC-2000;
                                                                                                                                                                                                                                     26-JUL-2001
                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                           leukaemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            166
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                                                                                                                                                                                                                                                                                                                                                                                                                                 25
                                                                              QA,
                          as
                                                     2001-442253/47
DB; AAI59623.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          eteklkeelsassartqnlkadlqrkeedyadlkekltdakkqikqvqkevsvmr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           VADKLRKDLFSSRSKLQTVYSELDQAKLELKSAQKDLQSADKEIMSLKKKLTMLQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ---VVSLQQALGKAEMLCSTLKKQMKYLEQQQDETKQAQEEAGRLRSKMKTMEQIELLLQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   eeleqqieklqaevkgykdennrlkekehknqddllkeketliqqlkeelqeknvtldvq 1098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EFLKNELDNVRAQL-SQKD-----KEK-----RDSQVIIDTLRDTLEERNAT----
                                                                                                YT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             rnlkefqehlqd--
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SQLPEVEEMIRDMGVGQSAVEQLAVYCVSLKKEYENL------KEARKASGE
                                  nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        50;
                          central nervous
                                                                                                                 HYSEQ INC
                                                                              Liu C
Wang Z
Zhou
                                                                                                                                                                                                                                                                                                                                                                                                               standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1780
                                                                                                                                  2000US-0488725.
2000US-0553317.
2000US-0558042.
2000US-0623312.
2000US-0653450.
2000US-0653450.
2000US-0653936.
2000US-0727344.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                           u C,
1 Z,
1 P,
                                                                                                                                                                                                                   2000WO-US34263
                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                  acids
                                                                                                                                                                                                                                                                                                   thrombolytic; drug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AA
                                                                                                                                                                                                                                                                                                                                                         SEQ
                                                                                        Asundi V,
Wehrman T,
                                                                              Goodrich
                                                                                                                                                                                                                                                                                                                                                                                                               Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -svkntkdlnvkelklkeeitqltnnlqdmkhllqlkeeeeetnrq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14.2%;
                                                                                                                                                                                                                                                                                                                                                         ID NO
                        polypeptides, system injuri
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                                                                                                                                                                                                                                                                                                                                                         5398
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                                                                              ₽,
                                                                                                Chen
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                                                                                        Χ'n
                         injuries
                                                                             len R, Ma
lu C, Xue
Drmanac R
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                                                                                                                                                                                                                                                                                                   screening;
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                                  useful
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RT;
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                                                                                       Qian XB,
Yang Y,
                                   for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        84;
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                                  treating
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                                                                                       Ren F,
Zhang
                                   disorders
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Example

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RESULT 26
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to human nucleic acids (AAI57798-AAI61369) and the encoded polypeptides (AAM38642-AAM42213) with nootropic, immunosuppressant and cytostatic activity. The polynucleotides are useful in gene therapy. A composition containing a polypeptide or polynucleotide of the invention may be used to treat diseases of the peripheral nervous system, such as peripheral nervous injuries, peripheral neuropathy and localised neuropathies and central nervous system diseases, such as localised neuropathies and central nervous system disease, such as Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager Syndrome. Other uses include the utilisation of the activity, chemotactic/chemokinetic activity, haemostatic and thrombolytic activity, cancer diagnosis and therapy, drug screening, and the polyper syndrome. The control of the activity, cancer diagnosis and therapy, drug screening, and the polyper syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1167
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C.N.S disorders.
Note: The sequence data for this
                 Region
                                            Region
                                                                       Region
                                                                                                    Region
                                                                                                                               Region
                                                                                                                                                            Protein
                                                                                                                                                                                                                     Peptide
                                                                                                                                                                                                                                   Key
                                                                                                                                                                                                                                                              Streptococcus
                                                                                                                                                                                                                                                                                          diagnosis
                                                                                                                                                                                                                                                                                                     Bovine lactoferrin binding protein;
                                                                                                                                                                                                                                                                                                                                 Streptococcus uberis bovine lactoferrin binding protein;
                                                                                                                                                                                                                                                                                                                                                                26-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                             AAW63043;
                                                                                                                                                                                                                                                                                                                                                                                                                      AAW63043 standard; Protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22 EFLKNELDNVRAQL-SQKD-----KEK-----RDSQVIIDTLRDTLEERNAT----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ---VVSLQQALGKAEMLCSTLKKQMKYLEQQQDETKQAQEEAGRLRSKMKTMEQIELLLQ 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       VADKLRKDLFSSRSKLQTVYSELDQAKLELKSAQKDLQSADKEIMSLKKKLTMLQ 220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  rnlkefqehlqd---svkntkdlnvkelklkeeitqltnnlqdmkhllqlkeeeeetnrq
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 eeleqqieklqaevkgykdennrlkekehknqddllkeketliqqlkeelqeknvtldvq 1106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SQLPEVEEMIRDMGVGQSAVEQLAVYCVSLKKEYENL------KEARKASGE 165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             50;
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                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                            uberis strain su-1 (ATCC
                                            /note=
282..32
                /note=
326..3:
                                                                                                                                                           52..561
                                                                         213..271
                                                                                                     200..212
                                                                                                                                148..199
                                                                                                                                                                                       /note=
                                                                                                                                                                                                /label=_Sig_peptide
                                                                                                                                                                                                                                Location/Qualifiers
                                                                                     /note=
                                                                                                                 /note-
                                                                                                                                           /label=_Mat_protein
                                              . . 325
                 . . 339
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21.3%;
                                                          "central
                                                                                                                                                                                       "alternative translation start site
                            "central repeated amino
                                                                                      "central
                                                                                                                "central repeated amino
 "central
                                                                                                                                                                       Met-11"
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                                                                                                                                                                                                                                                                                                                                                                                                                        561
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 151.5; DB 22; Length Pred. No. 0.00075; 62; Mismatches 84; Indels
 repeated
                                                        repeated
                                                                                     repeated
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                                                                                                                                                                                                                                                                                                      LBP;
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 amino
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                                                                                     amino
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 acid
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sequence B2"
                             sequence
                                                        sequence
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                               A2 "
                                                          Cl.
                                                                                     B1 "
                                                                                                                 A1 "
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RESULT 27
AAB21227
ID AAB212
XX
AC AAB212
XX
AC AAB212
XX
DT 09-MAR
XX
DE Protei
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This is the bovine lactoferrin binding protein (LBP) of Streptococcus uberis su-1. Its amino acid sequence was deduced from the novel isolated lbp gene (see AN42601). The LBP is lactoferrin species-specific; human lactoferrin does not effectively block binding of bovine lactoferrin. The invention provides recombinant vectors, transformed host cells and methods of producing recombinant bovine LBP of S. uberis. The bovine LBP, immunogenic fragments and/or chimeric proteins can be used, either alone or in combination with other antigens, in novel subunit vaccines for the prevention and treatment of S. uberis infections, particularly mastitis, as well as in diagnostic methods for determining the accessor of s.
                                                                                                      AAB21227 standard; Protein; 721
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunogenic Streptococcus uberis protein(s) that bind bovine lactoferrin - associated regulatory protein, useful in vaccin treatment and prevention of mastitis
               Protein encoded by tobacco NtMFP1-1 cDNA.
                                            09-MAR-2001 (first entry)
                                                                          AAB21227;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         determining the presence of S. uberis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-NOV-1996;
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                                                                                                                                                                                                                                         370
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                                                                                                                                                                                                                                                                                                                                                                                      10 DLAQEEENVLDREFLKN-----ELDNVRAQLSQKDKEKRDSQVIIDTLRDTLEERNATV 63
                                                                                                                                                                                                                                                                                                                            64 VSLQQALGKAEMLCSTLKKQMKYLEQQQDETKQAQEEAGRLRSKMKTMEQIELLLQSQLP 123
                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                 -----LFSSRSK-LQTVYSELDQAKLELKSAQKDLQSADKEIMSLKKKL 216
                                                                                                                                                                                                                                       eleeqveklveeitaakkeaeelqakaeglekdfeavkaekealeaeia-klkedhqkev
                                                                                                                                                                                                                                                                     EVEE----MIRDMGVGQSAVEQLAVYCVSLKKEYENLKEARKA-SGEVADKLRKD-----
                                                                                                                                                                                                                                                                                                  ddlqkkvmkaeqemkelsaqleeakeeletekaklaesekenaklteerdaakkeaekvp
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                                                                                                                                                                                dalnalladkekmlknlqdqldkakee---amkneqmsqeekaklqael
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Maclachlan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               561 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                      53;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 151; DB 19;
Pred. No. 0.0002;
                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                       3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           infections
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 561;
                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                474
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                                                                                                                                                                                                                                                                                                                                                                                                                      26;
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RESULT 2
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Best Local S
Matches 55
                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence is encoded by NtMFPI-1 cDNA from tobacco. It is a homologue of the matrix attachment region (MAR) binding filament-like protein 1 (MFPI) from tomato. MFPI has features of a novel anchor protein that most likely connects chromatin via MAR DNA with the nuclear envelope and nuclear filament proteins. MFPI nucleic acids and proteins may be used to better understand the mechanisms underlying this process so that the attachment of transgenes to the nuclear matrix may be used routinely to improve gene expression. They may be used to study MFPI expression, leading to the creation of novel developmental phenotypes that may be beneficial for crop growth and development. In addition, if the reduction in expression of one of the genes leads to a growth or developmental defect in the plant, this gene can be used as a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acid fragments from tobacco, corn, soybean and rice, encoding proteins that are homologs to the MAR binding filament-like protein 1 (MFP1), useful for development of novel phenotypes -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; Page 45-47;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Harder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200061615-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Misc-difference
   AAM78520 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (DUPO ) DU PONT DE NEMOURS & CO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tobacco; MAR-binding filament-like protein 1; MFP1;
matrix attachment region; MAR; anchor protein.
                                                                                                                                                                                                                                                                                                                                                                                                                    novel herbicide target.
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                                                                                                                                                 LLQSQLPEVEEMIRDMGVGQSAVEQLAVYCVSLKKEYENLKEARKASGEVADKLRKDLF-
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                                                                                                                                                                                                                        SLQQALGKAEMLCSTLKKQMKYLEQQQDETK-----QAQEEAGRLRSKMKTMEQIEL
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                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                     SSRSKLQTVYSELDQAKLELKSAQKDLQSADKEIMSLKKKLTMLQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           62pp; English.
                                                                                                                                                                                                                                                                                                                              14.2%;
23.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "encoded by GAGATT"
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                                                                                                                                                                                                                                                                                                                 46;
   990
                                                                                                                                                                                                                                                                                                              Score 151; DB 21;
Pred. No. 0.00027;
6; Mismatches 99,
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Matches
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Best Local
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27-APR-2000;
20-JUN-2000;
19-JUL-2000;
01-SEP-2000;
15-SEP-2000;
20-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                     The invention relates to polynucleotides (AAK51456-AAK53435) and the encoded polypeptides (AAM78323-AAM80302) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polynucleotides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides have various cytokine-like activities, e.g. stem cell growth factor activity, haematopoiesis regulating activity, tissue growth factor activity, immunomodulatory activity and activity in activity and may be useful in the diagnosis and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; cytokine; cell proliferation; cell differentiation; gene therapy; vaccine; peptide therapy; stem cell growth factor; haematopolesis; tissue growth factor; immunomodulatory.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 20; Page 3425-3427; 6221pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-476283/51.
N-PSDB; AAK51653.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tang YT,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nervous system disorder; arthritis; inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-NOV-2001
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                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic
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                                                                                                                                                                                                                                                                                were missing at the time of publication
                                                                                                                                                                                                                                                                                                                      Note:
                                                                                                                                                                                                                                                                                                                                      inflammation.
                                                                                                                                                                                                                                                                                                                                                        treatment of cancer, leukaemia, nervous system disorders, arthritis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HYSE-)
                                                                                                                                                                                                                                                                                             Note: Records for SEQ ID NO 2110 (AAK52581), (AAM80020) are omitted as the relevant pages
 773
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                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               QA,
                                  LGKAEMLCSTLKKQM-----KYLEQQQDETKQAQEEAGRLRSKMKTMEQIELLLQSQL
                                                                    enkelesekeqlkkglellkasfkkterlevsyggldiengrlgktlensnkkigglese
lqdlemenqtlqknleelkisskrleqlekenksleqetsqlekdkkqlekenkrlrqqa
                                                                                                          QEEENVLDREFLKNELDNVRAQLSQKDKEKRDSQ---VIIDTLRDTLEERNATVVSLQQA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   diagnosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2000US-0496914.

2000US-0560875.

2000US-0598075.

2000US-0598075.

2000US-0620325.

2000US-062325.

2000US-063561.

2000US-0693325.

2000US-0728422.
                                                                                                                                                   Conservative
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24.68;
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J, Zhang J, Ren F
man T, Goodrich R;
                                                                                                                                                 49;
                                                                                                                                               Score 150; DB 22;
Pred. No. 0.00048;
9; Mismatches 95;
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from
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Best Local S
Matches 61
                                                                                                     1482
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The specification describes laminin 12 which includes an alpha 2, beta and gamma 3 subunit. Laminin is a connective tissue adhesion molecule Laminin is useful for promoting tissue repair due to wounds and to promote nerve growth or regeneration. The present sequence represents human laminin beta 4.
1542 sneeadgaqkllvkakaaekaanillnldktlnqlqqaqitqgranstitqltanitkik 1601
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                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
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                                                QAQEEAG---RLRSKMKTMEQ----IELLLQSQLPEVEEMIRDMGVGQSAVEQLAVYCVSLK
                                                                                              {\tt ikkvknflleenvppediekvangvldihlpipsqnltdelvkiqkhmqlcedyrtdenr}
                                                                                                                                                 IDTLRDTLEERNATVVSLQQALG -----KAEMLCSTLKKQMKYL----EQQQDETK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   beta 1; gamma 3; subunit; nerve regeneration;
hesion; tissue repair; wound; nerve growth;
                                                                                                                                                                                                                                                                                                                        14.0%;
23.8%;
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                                                                                                                                                                                                                                                                                            Score 149.5; DB 20;
Pred. No. 0.0011;
Pred. No. 0.0011;
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RESULT 3
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and pharmaceutical compositions comprising a HUMAP protein, a HUMAP agonist or HUMAP antagonist. The HUMAPs acts as modulators of cell signalling, differentiation and proliferation. A HUMAP is useful for screening a compound for effectiveness as an agonist or antagonist of HUMAP activity. The protein, or the identified agonist or antagonist is useful for treating a disease or condition associated with decreased or increased expression of functional HUMAP. A HUMAP nucleic acid is useful for screening a compound for its ability to alter expression of that particular HUMAP gene. A wide variety of disease may be treated using compositions of the invention. These diseases include cell proliferative disorders (e.g., actinic keratosis, arteriosclerosis); cancer (e.g.,
                                                                                                                                                                                                                                                                                                                        The invention relates to 17 human membrane-associated proteins, HUMAP-1 to HUMAP-17 (AAB29644-B29660) and the cDNAs encoding them (AAC64274-C64290). The invention also relates to expression constructs, host cells and transgenic organisms comprising a HUMAP nucleic acid sequence; the recombinant preparation of a HUMAP; methods of screening compounds for their ability to modulate HUMAP activity or expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human membrane-associated protein, useful for diagnosis and treatment of cell signaling, cell differentiation and cell proliferation disorders such as cancer, and for identifying agonists and antagonists
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human membrane-associated protein: HUMAP: transgenic organism; drug screening; cell signalling modulator; agonist; antagonist; cell differentiation modulator; cell proliferation modulator; cell proliferative disorder; cancer; cell differentiation disorder; cell signalling disorders; endocrine disorder; developmental disorder; cell signalling disorders; endocrine disorder; hyperpituitarism; hypothyroidism; hypot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; Page 86-87; 99pp; English.
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23-JUN-1999;
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99US-0140580.
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Cd differentiation disorders, in particular developmental disorders (e.g.,
Cc renal tubular acidosis, anaemia, Cushing's syndrome, achondroplasia,
CC epilepsy, and muscular dystrophy); cell signalling disorders in
CC particular endocrine disorders such as hypothalmus and pituitary
CC disorders resulting from lessions such as thrombosis; disorders
CC associated with hyperpituitarism (e.g., acromegaly); disorders associated
CC with hypothyroidism (e.g., goitre); hyperparathyroidism; pancreatic
CC disorders such as type I or type II diabetes mellitus; infections;
CC immunological disorders; hereditary neuropathies (e.g.,
CC neurofibromatosis); and disorders associated with gonadal steroid
CC neurofibromatosis); and disorders associated with gonadal steroid
CC osteoporosis, Leydig cell deficiency and gynecomastia). Antibodies which
CC specifically bind HUMAP may be used for the diagnosis of disorders
CC associated with the expression of HUMAP, or in assays to monitor patients
CC being treated with HUMAP or agonists, antagonists or inhibitors of HUMAP.

The present sequence represents a HUMAP of the invention.
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Best Local
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03-FEB-2000;
27-APR-2000;
20-JUN-2000;
19-JUL-2000;
                                                                                                                                                                                          Homo
                                                                                                                                                                                                                        Human; cytokine; cell proliferation; cell differentiation; gene th vaccine; peptide therapy; stem cell growth factor; haematopoiesis; tissue growth factor; humanomodulatory; cancer; leukaemia; nervous system disorder; arthritis; inflammation.
                                                                                    05-FEB-2001;
                                                                                                                     09-AUG-2001
                                                                                                                                                     WO200157190-A2
                                                                                                                                                                                                                                                                                                             Human protein SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                    AAM78880 standard; Protein;
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2000US-0560875.
2000US-0598075.
2000US-0620325.
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Pred. No. 0.00017;
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                                                                                                                                                                                                                                                                         cell differentiation; gene therapy;
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Best Local S
Matches 67
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15-SEP-2000;
20-OCT-2000;
30-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to polynucleotides (AAK51456-AAK53435) and the encoded polypeptides (AAM78323-AAM80302) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polynucleotides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides have various cytokine-like activities, e.g. stem cell growth factor activity, haematopolesis regulating activity, tissue growth factor activity, hammatopolesis regulating activity, tissue growth factor activity, tinductory activity and activiny inhibin activity and may be useful in the diagnosis and/or treatment of cancer, leukaemia, nervous system disorders, arthritis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Note: Records for SEQ ID NO 2110 (AAK52581), 2111 (AAM80020) are omitted as the relevant pages from were missing at the time of publication.
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Zhao
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AAM79864;
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                        standard;
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Wang D,
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Wejhrman T,
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Wang J, Zhang J, Ren
Wejhrman T, Goodrich R;
                          Protein;
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d gene therapy -
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Pred. No. 0.00062;
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F, (
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06-NOV-2001

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Best Local Similarity 21.5
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27-APR-2000;
20-JUN-2000;
19-JUL-2000;
01-SEP-2000;
15-SEP-2000;
20-OCT-2000;
                                                                                                                                                                                                                                                                                 The invention relates to polynucleotides (AAK51456-AAK53435) and the encoded polypeptides (AAM78323-AAM80302) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polynucleotides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides have various cytokine-like activities, e.g. stem cell growth factor activity, haematopoiesis regulating activity, tissue growth factor activity, haematopoiesis regulating activity, tissue growth factor activity, the diagnosis and/or treatment of cancer, leukaemia, nervous system disorders, arthritis and
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Zhao
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                                                                                                                                                                                                                                    Note: Records for SEQ ID NO 2110 (AAK52581), (AAM80020) are omitted as the relevant pages were missing at the time of publication.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acids encoding polypeptides with cytokine-like useful in diagnosis and gene therapy - \,
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AJ,
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                                                keevkevlqaleelavnydqksqevedktkeyellsdelnqksatlasidaelqklkemt
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2000US-0560875
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, Zhang J, Ren
n T, Goodrich R;
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Pred. No. 0.00063;
5; Mismatches 94
                       -KQMKYLEQQQDETKQAQEEAGRLRSKMKTM 111
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1 F, Chen
                                                                                                                                                                         DB 22;
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Best Local S
Matches 57
 1302
                                                                                                                                                                        The invention relates to an isolated nucleic acid detection reagent capable of detecting 1000 or more genes from Drosophila. The invention useful in developmental biology and in elucidating cell signalling and cell-cell interactions in higher eukaryotes for the development of insecticides, therapeutics and pharmaceutical drugs. The invention discloses genomic DNA sequences (ABL16176-ABL30511), expressed DNA sequences (ABL01840-ABL16175) and the encoded proteins
                                                                                                                                                                                                                                                                                                                                                                  WPI; 200
N-PSDB;
                                                                                                                                     The sequence data for this patent did not form specification, but was obtained in electronic at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                 New isolated nucleic acid genes from Drosophila and interactions .
                                                                                                                                                                                                                                                                                       Disclosure; SEQ ID NO 4824; 21pp + Sequence Listing; English
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11-JUL-2000;
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DB; ABL03447.
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                                                 57; Conservative
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Pred. No. 0.0019;
9; Mismatches 102;
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          CC polypeptide (II) sequences. (I) is useful as hybridisation probes, CC polymerase chain reaction (PCR) primers, oligomers, and for chromosome CC and gene mapping, and in recombinant production of (II). The CC and gene mapping, and in recombinant production of (II). The CC constant fying expressed genes. (I) is useful in gene therapy techniques CC for identifying expressed genes. (I) is useful in gene therapy techniques CC (II), (II) is useful for generating antibodies against it, detecting or CC quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical cimaging of sites expressing (II). (I) and (II) are useful for treating CC disorders involving aberrant protein expression or biological activity. The polypeptide and polynucleotide sequences have applications in CC diagnostics, forensics, gene mapping, identification of mutations cresponsible for genetic disorders or other traits to assess biodiversity amino acid sequences. ABG00010-ABG3037 represent novel human
                                                                                                                                                                                                                                                                                                                                                       New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess
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food supplement; r
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                                                                                                                                                                                                                                                                           The invention relates to isolated polynucleotide (I) and
                                                                                                                                                                                                                                                                                                              Claim 20;
                                                                                                                                                                                                                                                                                                                                              biodiversity
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23-AUG-2000;
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2000US-0649167
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mapping; gene mapping; gene therapy; forensic medical imaging; diagnostic; genetic disorder.
   acid
                                                                                                                                                                                                                                                                                                           No 36864; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tang
   sequences of the invention
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Tang Wang Zhao

Wang Z, Zhou P,

Asundi V, Wehrman T,

Chen R,

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WPI; 2001-442253/47. N-PSDB; AAI58253.

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                                                                                                                                                                          19-JUL-2000;
03-AUG-2000;
                                                                                                                                                                                                 25-APR-2000;
09-JUL-2000;
                                                                                                                                                                                                                                                                                               WO200153312-A1
                                                                                                                                                                                                                                                                                                                                                         chemokinetic; thrombolytic; drug screening; arthritis; inflammation;
                                                                                                                                                                                                                                                                                                                                                                    Alzheimer's; Parkinson's disease; Huntington's disease; haemosta amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic;
                                                                                                                                                                                                                                                                                                                                                                                                      Human; nootropic;
                                                                                                                                                                                                                                                                                                                                                                                                                              Human polypeptide SEQ ID NO 2242
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29-NOV-2000;
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                                                                                                                                                                                                                                                                                                                                                                                           peripheral
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57; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard;
                                                                                                                                                                                                                                                                                                                                                                                           nervous
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2000US-0553317.

2000US-0598042.

2000US-0620312.

2000US-0653450.

2000US-0663450.

2000US-0693036.

2000US-0727344.
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                                                                                                                                                                                                                                                2000WO-US34263
                                                                                                                                                                                                                                                                                                                                                                            immunosuppressant; cytostatic; gene therapy; cance
s system; neuropathy; central nervous system; CNS;
inson's disease; Huntington's disease; haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Protein;
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25.1%;
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AC AAM4
AC AAM4
AC BE Huma
XX
BE Huma
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KW Huma
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COS Homc
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 21-JAN-2000;
25-APR-2000;
09-JUL-2000;
19-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to human nucleic acids (AAI57798-AAI61369) and the encoded polypeptides (AAM38642-AAM42213) with nootropic, immunosuppressant and cytostatic activity. The polynucleotides are useful in gene therapy. A composition containing a polypeptide or polynucleotide of the invention may be used to treat diseases of the peripheral nervous system, such as peripheral nervous injuries, peripheral neuropathy and localised neuropathies and central nervous system diseases, such as Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager Syndrome. Other uses include the utilisation of the activity prager Syndrome. Other uses include the utilisation of the activity, chemotactic/chemokinetic activity, haemostatic and thrombolytic activity, cancer diagnosis and therapy, drug screening, assays for receptor activity, arthritis and inflammation, leukaemias and C.N.S discrete
                                                                                                                                                                                                                       Human; nootropic; immunosuppressant; cytostatic; gene therapy; cancer; peripheral nervous system; neuropathy; central nervous system; CNS; Alzheimer's; Parkinson's disease; Huntington's disease; haemostatic; amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic; chemokinetic; thrombolytic; drug screening; arthritis; inflammation;
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                                                                                                               26-JUL-2001.
                                                                                                                                               WO200153312-A1
                                                                                                                                                                             Homo sapiens.
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2000US-0488725.
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2000US-0598042.
2000US-0620312.
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14-SEP-2000;
19-OCT-2000;
29-NOV-2000;
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Wang
                                                                              18-OCT-2000
                                                                                                        AAG46982;
                                                                                                                                   AAG46982 standard;
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                                                                                                                                                                                                 DLFSSRSKLQTVYSELDQAKLELKSAQKDLQSADK----EIMSLKKKL
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Xu C,
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Protein identification; signal transduction pathway; metabolic pathway; hybridisation assay; genetic mapping; gene expression control; promoter; termination sequence.

Arabidopsis

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protein

fragment SEQ ID NO:

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Best Local
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                                                                                                                                                                                                                                                             The invention relates to polynucleotides (AAK51456-AAK53435) and the encoded polypeptides (AAM78323-AAM80302) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polynucleotides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides have various cytokine-like activities, e.g. stem cell growth factor activity, haematopoiesis regulating activity, tissue growth factor activity, immunomodulatory activity and activiny/inhibin activity and may be useful in the diagnosis and/or treatment of cancer, leukaemia, nervous system disorders, arthritis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-FEB-2000;
27-APR-2000;
20-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic acids encoding polypeptides with cytokine-like useful in diagnosis and gene therapy -
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15-SEP-2000;
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Wang D,
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  Score 146; DB 22;
Pred. No. 0.00095;
50; Mismatches 95;
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The invention relates to isolated polynucleotide (I) and polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polynucleotides are also used in diagnostics as expressed sequence tags for identifying expressed genes. (I) is useful in gene therapy techniques to restore normal activity of (II) or to treat disease states involving (II). (II) is useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful for treating disorders involving aberrant protein protein are useful for treating disorders involving aberrant protein sequences have applications in the polypeptide and polypucleotide sequences have applications in
                                                                                                                                                                                                                                                                               New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutation responsible for genetic disorders or other traits and to assess biodiversity
                                                                                                                                                                                                                                            Claim 20; SEQ ID
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23-AUG-2000;
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2000US-0649167.
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                                                                                                                                                                                                                                                                                                   responsible for genetic disorders or other traits to assess biodiversity and to produce other types of data and products dependent on DNA and amino acid sequences. ABG00010-ABG30377 represent novel human diagnostic amino acid sequences of the invention.

Note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                     185 YSELDQAKLELKSAQKDLQSADKEIM-SLKKKLTML 219
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| 263 lqeenrqkls1stklkqleeeeeeakhnlekqiatl 1298
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